

## 5-Chloro-2-nitroaniline

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## Key indicators

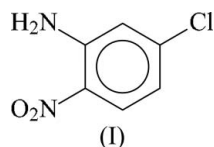
Single-crystal X-ray study  
 $T = 295$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.002$  Å  
 $R$  factor = 0.041  
 $wR$  factor = 0.122  
Data-to-parameter ratio = 13.4For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.The title compound,  $\text{C}_6\text{H}_5\text{ClN}_2\text{O}_2$ , exists as a planar molecule; adjacent molecules are linked by  $\text{N} \cdots \text{O}$  hydrogen bonds into ribbons.

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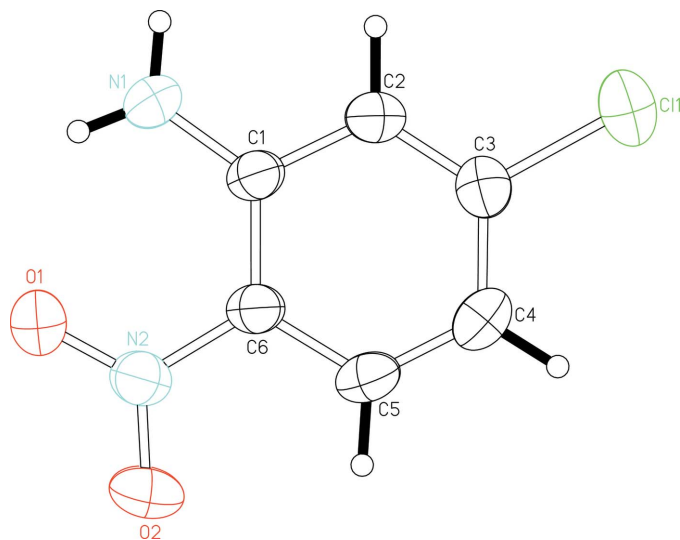
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## Comment

Chloronitroanilines possess non-linear optical properties. The crystal structure of 5-chloro-2-nitroaniline, (I), was presented at a conference (Gerbi *et al.*, 1988), but it has not been abstracted into the Cambridge Structural Database (Version 5.26; Allen, 2002). The first chloronitroaniline to be authenticated was 2-chloro-4-nitroaniline (McPhail & Sim, 1965); chloronitroanilines are generally not well studied, and 4,5-dichloro-2-nitroaniline (Doyle, 1999) represents a rare example. The title compound (Fig. 1) exists as a planar molecule; adjacent molecules are linked by hydrogen bonds (Table 1) into a ribbon (Fig. 2).

## Experimental

The title compound was obtained commercially and was recrystallized from dimethyl sulfoxide.



**Figure 1**  
ORTEP plot (Johnson, 1976) of (I), with displacement ellipsoids drawn at the 50% probability level. H atoms are drawn as spheres of arbitrary radii.

Crystal data

C<sub>6</sub>H<sub>5</sub>ClN<sub>2</sub>O<sub>2</sub>  
*M<sub>r</sub>* = 172.57  
 Triclinic, *P* $\bar{1}$   
*a* = 7.073 (3) Å  
*b* = 7.423 (3) Å  
*c* = 7.711 (3) Å  
 $\alpha$  = 83.87 (3)°  
 $\beta$  = 81.98 (3)°  
 $\gamma$  = 62.24 (3)°  
*V* = 354.4 (3) Å<sup>3</sup>

*Z* = 2  
*D<sub>x</sub>* = 1.617 Mg m<sup>-3</sup>  
 Mo *K*α radiation  
 Cell parameters from 3038 reflections  
 $\theta$  = 3.1–27.5°  
 $\mu$  = 0.48 mm<sup>-1</sup>  
*T* = 295 (2) K  
 Plate, yellow  
 0.32 × 0.21 × 0.01 mm

Data collection

Rigaki R-AXIS RAPID IP diffractometer  
 $\omega$  scans  
 Absorption correction: multi-scan (ABSCOR; Higashi, 1995)  
*T<sub>min</sub>* = 0.505, *T<sub>max</sub>* = 0.955  
 3525 measured reflections

1608 independent reflections  
 1299 reflections with *I* > 2σ(*I*)  
*R<sub>int</sub>* = 0.023  
 $\theta_{max}$  = 27.5°  
*h* = -9 → 9  
*k* = -7 → 9  
*l* = -9 → 9

Refinement

Refinement on *F*<sup>2</sup>  
*R* [*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.041  
*wR* (*F*<sup>2</sup>) = 0.122  
*S* = 1.07  
 1608 reflections  
 120 parameters

All H-atom parameters refined  
 $w = 1/[\sigma^2(F_o^2) + (0.0843P)^2]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 ( $\Delta/\sigma$ )<sub>max</sub> = 0.001  
 $\Delta\rho_{max}$  = 0.38 e Å<sup>-3</sup>  
 $\Delta\rho_{min}$  = -0.21 e Å<sup>-3</sup>

Table 1

Hydrogen-bond geometry (Å, °).

<i>D</i> —H··· <i>A</i>	<i>D</i> —H	H··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> —H··· <i>A</i>
N1—H11···O1	0.84 (1)	2.07 (2)	2.633 (2)	124 (2)
N1—H11···O1 <sup>i</sup>	0.84 (1)	2.45 (2)	3.152 (2)	141 (2)
N1—H12···O2 <sup>ii</sup>	0.85 (1)	2.35 (1)	3.113 (2)	150 (2)

Symmetry codes: (i) -*x* + 1, -*y*, -*z* + 2; (ii) *x* + 1, *y*, *z*.

H atoms were located in difference Fourier maps and were refined with distance restraints of N—H = 0.85 (1) Å and C—H = 0.95 (1) Å.

Data collection: *RAPID-AUTO* (Rigaku, 1998); cell refinement: *RAPID-AUTO*; data reduction: *CrystalStructure* (Rigaku/MS, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

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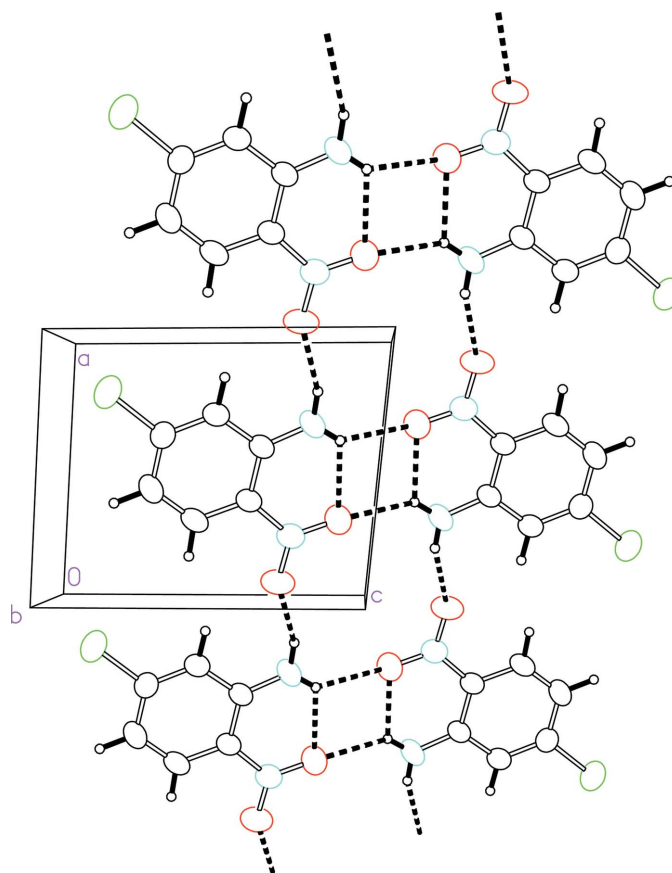


Figure 2  
 ORTEPII plot (Johnson, 1976) of the ribbon structure of (I). Dashed lines indicate hydrogen bonds.

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